

Notice of References Cited	Application/Control No. 10/762,154	Applicant(s)/Patent Under Reexamination NEZU ET AL.	
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U.S. PATENT DOCUMENTS

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	U	Skolnick et al. From genes to protein structure and function: novel applications of computational approaches in the genomic era. Trends in Biotech 18(1): 34-39, 2000. ✓
	V	Bork, A. Powers and pitfalls in sequence analysis: the 70% hurdle. Genome Res 10: 398-400, 2000. ✓
	W	Doerks et al. Protein annotation: detective work for function prediction. Trends in Genetics 14(6): 248-250, 1998. ✓
	X	Smith et al. The challenges of genome sequence annotation or "The devil is in the details". Nature Biotech 15: 1222-1223, 1997. ✓

*A copy of this reference is not being furnished with this Office action. (See MPEP § 707.05(a).)
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	X	Ngo et al. Computational complexity, protein structure prediction, and the Levinthal paradox. The Protein Folding Problem and Tertiary Structure Prediction, pp. 492-495, 1994. ✓

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